

- PI 591445. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1148; UCDS/P-A(H)2. Pedigree - UCD S/P-A(H)2.  
Germplasm pool selected from CUF101 for increased number of seeds per  
pod through 2 cycles of phenotypic recurrent selection. Approx. 45  
plants intercrossed in each cycle.
- PI 591446. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1280; UCDS/P-B(H)2. Pedigree - UCD S/P-B(H)2.  
Germplasm pool selected from Team for increased number of seeds per pod  
through 2 cycles of phenotypic recurrent selection. Approx. 45 plants  
intercrossed in each cycle.
- PI 591447. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1167; UCWSS-B(L)2. Pedigree - UC WSS-B(L)2.  
Germplasm pool selected from Team for decreased seed yield through 2  
cycles of phenotypic recurrent selection. Approx. 45 plants intercrossed  
in each cycle.
- PI 591448. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1168b; UCWSS-B(H)2. Pedigree - UC WSS-B(H)2.  
Germplasm pool selected from Team for increased seed yield through 2  
cycles of phenotypic recurrent selection. Approx. 45 plants intercrossed  
in each cycle.
- PI 591449. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1175; UCWSS-C(H)2. Pedigree - UC WSS-C(H)2.  
Germplasm pool selected from Vernal for increased seed yield through 2  
cycles of phenotypic recurrent selection. Approx. 45 plants intercrossed  
in each cycle.
- PI 591450. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1155; UCDS-B(H)2. Pedigree - UC DS-B(H)2.  
Germplasm pool selected from Team for increased seed yield through 2  
cycles of phenotypic recurrent selection. Approx. 45 plants intercrossed  
in each cycle.
- PI 591451. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1161; UCDS-C(H)2. Pedigree - UC DS-C(H)2.  
Germplasm pool selected from Vernal for increased seed yield through 2  
cycles of phenotypic recurrent selection. Approx. 45 plants intercrossed  
in each cycle.

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- PI 591452. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1269; UCNE-A(L)2. GP-281. Pedigree -  
UCNE-A(L)2. Germplasm pool selected from CUF101 for decreased nectar  
production through 2 cycles of phenotypic recurrent selection. Approx.  
45 plants intercrossed in each cycle.
- PI 591453. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1270; UCNE-A(H)2. GP-282. Pedigree - UC  
NE-A(H)2. Germplasm pool selected from CUF101 for increased nectar  
production through 2 cycles of phenotypic recurrent selection. Approx.  
45 plants intercrossed in each cycle of selection.
- PI 591454. *Medicago sativa* L. ssp. *sativa*  
Breeding. Population. UC-1272; UCNE-B(L)2. GP-283. Pedigree - UC  
NE-B(L)2. Germplasm pool selected from Team for decreased nectar  
production through 2 cycles of phenotypic recurrent selection. Approx.